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RAW SEQUENCE LISTING DATE: 03/04/2002
PATENT APPLICATION: US/09/983,536 TIME: 08:35:16

Input Set : A:\620-173.app
Output Set: N:\CRF3\03042002\I983536.raw

<110> APPLICANT: Carrillo, Nestor J
Valle, Estela M
Tognetti, Vanesa B
Palatnik, Javier F
Castejon, Maria FF
<120> TITLE OF INVENTION: Stress Tolerant Plants
<130> FILE REFERENCE: 620-173
<140> CURRENT APPLICATION NUMBER: US 09/983,536
<141> CURRENT FILING DATE: 2001-10-24
<160> NUMBER OF SEQ ID NOS: 6
<170> SOFTWARE: PatentIn Ver. 2.1
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<211> LENGTH: 706
<212> TYPE: DNA
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Nucleic acid molecule encoding fusion polypeptide
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tgtcttcaaa aaggtttcat tgaacaatgt ttctataagt ggaagggtag gcaccatcag 180
agctctcata atgtcaaaga aaattggttt attctacggt actcaaactg gtaaaactga 240
atcagtagca gaaatcattc gagacgagtt tggtaatgat gtggtgacat tacacgatgt 300
ttcccaggca gaagtaactg acttgaatga ttatcaatat ttgattattg gctgtcctac 360
ttggaatatt ggcgaactgc aaagcgattg ggaaggactc tattcagaac tggatgatgt 420
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taattttcag gatgcgatcg gtattttgga agaaaaaatt tctcaacgtg gtggtaaaac 540
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<210> SEQ ID NO: 2
<211> LENGTH: 233
<212> TYPE: PRT
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Predicted protein sequence of transit peptide and flavodoxin protein
<400> SEQUENCE: 2
Asp Pro Ser Ser Thr Thr Thr Thr Asn Met Ala Ala Ala Val Thr Ala
1 5 10 15
Ala Val Ser Leu Pro Tyr Ser Asn Ser Thr Ser Leu Pro Ile Arg Thr

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59              20                  25                  30
61 Ser Ile Val Ala Pro Glu Arg Leu Val Phe Lys Lys Val Ser Leu Asn
62          35                  40                  45
64 Asn Val Ser Ile Ser Gly Arg Val Gly Thr Ile Arg Ala Leu Ile Met
65      50                  55                  60
67 Ser Lys Lys Ile Gly Leu Phe Tyr Gly Thr Gln Thr Gly Leu Thr Glu
68  65                  70                  75                  80
70 Ser Val Ala Glu Ile Ile Arg Asp Glu Phe Gly Asn Asp Val Val Thr
71                  85                  90                  95
73 Leu His Asp Val Ser Gln Ala Glu Val Thr Asp Leu Asn Asp Tyr Gln
74              100                 105                 110
76 Tyr Leu Ile Ile Gly Cys Pro Thr Trp Asn Ile Gly Glu Leu Gln Ser
77          115                 120                 125
79 Asp Trp Glu Gly Leu Tyr Ser Glu Leu Asp Asp Val Asp Phe Asn Gly
80      130                 135                 140
82 Lys Leu Val Ala Tyr Phe Gly Thr Gly Asp Gln Ile Gly Tyr Ala Asp
83 145                 150                 155                 160
85 Asn Phe Gln Asp Ala Ile Gly Ile Leu Glu Glu Lys Ile Ser Gln Arg
86                  165                 170                 175
88 Gly Gly Lys Thr Val Gly Tyr Trp Ser Thr Asp Gly Tyr Asp Phe Asn
89              180                 185                 190
91 Asp Ser Lys Ala Leu Arg Asn Gly Lys Phe Val Gly Leu Ala Leu Asp
92          195                 200                 205
94 Glu Asp Asn Gln Ser Asp Leu Thr Asp Asp Arg Ile Lys Ser Trp Val
95      210                 215                 220
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98 225                 230
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102 <211> LENGTH: 162
103 <212> TYPE: DNA
104 <213> ORGANISM: Pisum sativum
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108 atcagaacat ctattgttgc accagagaga cttgtcttca aaaaggtttc attgaacaat 120
109 gtttctataa gtggaagggt aggcaccatc agagctctca ta                    162
112 <210> SEQ ID NO: 4
113 <211> LENGTH: 54
114 <212> TYPE: PRT
115 <213> ORGANISM: Pisum sativum
117 <400> SEQUENCE: 4
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119   1               5                  10                  15
121 Thr Ser Leu Pro Ile Arg Thr Ser Ile Val Ala Pro Glu Arg Leu Val
122             20                  25                  30
124 Phe Lys Lys Val Ser Leu Asn Asn Val Ser Ile Ser Gly Arg Val Gly
125          35                  40                  45
127 Thr Ile Arg Ala Leu Ile
128      50
131 <210> SEQ ID NO: 5
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Input Set : A:\620-173.app
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132 <211> LENGTH: 23
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134 <213> ORGANISM: Artificial Sequence
136 <220> FEATURE:
137 <223> OTHER INFORMATION: Description of Artificial Sequence: Primer
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145 <212> TYPE: DNA
146 <213> ORGANISM: Artificial Sequence
148 <220> FEATURE:
149 <223> OTHER INFORMATION: Description of Artificial Sequence: Primer
151 <400> SEQUENCE: 6
152 actgtcgact ttttacaaac caaat 25

VERIFICATION SUMMARY DATE: 03/04/2002
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